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Best Local Sim:
Matches 1321;
                                                                                                                                                                                                                                                                                                                                                            Receptor protein inducing differentiation extracellular region only and can be used
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Protein 7F4; differentiation; osteoblast cell; bone growth; bone sarcoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic acid
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                                                                                                                                                                                                                         The present sequence encodes a protein designated 7F4. This protein is capable of inducing differentiation in osteoblast cells. The protein may be used to screen compounds for the ability to bind to it, for use as ligands, agonists or antagonists and inhibiting or otherwise altering its differentiation inducing activity. Compounds so identified, as well as the protein itself, DNA encoding it, and antibodies to it, may be used in treatment of diseases of bone growth and osteoblast differentiation,
                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1998-568275/48.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Kimura N,
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                                                                                                                                                                                         Sequence
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                       TTCTGCAGGTACATTTGTCAAGGCGCCCTGCGAAATCCCCCCATACTCAAGGACAATGTGA
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                                                                                                                                                                                        1509 BP; 387 A; 403 C; 294 G; 425 T; 0 U; 0 Other;
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                                                                                                                                                                                                  Sequence 1509 BP; 387 A; 403 C; 294 G; 425 T; 0 U; 0 Other;
                                                                                                                                                                                                                                           The present invention relates to a transgenic non-human animal having the expression of 7F4 gene artificially modified. The transgenic animals are a disease model for bone and glycolipid metabolism disorders. Substances identified by the screening method are agents for the prevention and treatment of diseases including osteoporosis, obesity and diabetes. The present sequence represents the modified mouse 7F4 encoding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Transgenic non-human animal with modified expression of 7F4 gene screening remedies for bone or glycolipid metabolism disorders.
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P-PSDB; ADM46623.
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glycolipid metabolism disorder; osteoporosis; obesity; diabetes; ds.
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                                                                                                                           AGCATTCAAACACATGAGTCTATGAGGGCCAAACCAATTCAAACCACCACAGGTTAACAA
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Novel tmst2-receptor polypeptide useful for diagnosing and treating disease e.g., autoimmune disease, cachexia, cancer or viral, bacterial
                                                       WPI; 2004-224390/21.
N-PSDB; ADJ45749.
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Best Local Similarity
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                                                                                              Saris C;
                                                                                                                                             09-JUL-1999;
07-JUL-2000;
                                                                                                                                                                             17-JUL-2003; 2003US-00622407.
                                                                                                                                                                                                                                                                                acquired immunodeficiency syndrome; AIDS; anaemia; autoimmune disease; cachexia; cancer; cerebral malaria; diabetes mellitus; disseminated intravascular coagulation; haemorrhagic shock; hepatitis; insulin resistance; lgprosy; leukaemia; lymphoma; meningitis; multiple sclerosis; ischaemia; obesity; organ rejection; rheumatoid arthritis; septic shock; stroke; adult respiratory distress syndrome; ARDS; tuberculosis; viral disease
                                                                                                                       (SARI/)
                                                                                                                                                                                                            29-JAN-2004.
                                                                                                                                                                                                                                    US2004018544-A1.
                                                                                                                                                                                                                                                            Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                Mouse; tmst2-receptor; tmst2 00004-d1;
                                                                                                                                                                                                                                                                                                                                                                                                       Murine tmst2 00004-d1 polypeptide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADJ45750;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADJ45750 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to transmembrane decoy-receptor (tmst2) proteins and their secreted splice variants, belonging to the tumour necrosis factor (TNP) receptor super gene family and polynucleotides encoding such or ameliorating diseases associated with or resulting from abnormal tmsting and/or abnormal expression of its putative ligand, such as sepsis, and parasitic diseases or cancer. They may also be used for chromosome present sequence is mouse tmst2-receptor protein used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 133
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                                                                                                                    SARIS C.
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Best Local Similarity
Matches 133; Conserv
       Receptor protein inducing differentiation extracellular region only and can be used treatment of bone growth disorders.
                                                       WPI; 1998-568275/48.
N-PSDB; AAV68046.
                                                                                       Kimura N,
                                                                                                     (CHUG-) CHUGAI RES INST MOLECULAR MEDICINE INC
                                                                                                                                01-APR-1997;
                                                                                                                                                     01-APR-1998;
                                                                                                                                                                            08-OCT-1998.
                                                                                                                                                                                                 WO9843998-A1
                                                                                                                                                                                                                                Protein
                                                                                                                                                                                                                                                                                   Unidentified.
                                                                                                                                                                                                                                                     Peptide
                                                                                                                                                                                                                                                                                                       Protein 7F4;
                                                                                                                                                                                                                                                                                                                          Amino acid sequence of
                                                                                                                                                                                                                                                                                                                                                  28-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                       AAW80254;
                                                                                                                                                                                                                                                                                                                                                                                      AAW80254 standard; protein; 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a tmst2-receptor polypeptides and the polynucleotide encoding them. The sequences of the invention are useful for treating diseases and conditions including acquired immundeficiency syndrome (AIDS), anaemia, autoimmune diseases, cachexia, cancer, cerebral haemorrhagic shock, hepatitis, insulin resistance, leprosy, leukaemia, lymphoma, meningitis, multiple sclerosis, ischaemia, obesity, organ distress syndrome (ARDS), tuberculosis and a number of viral diseases. This sequence represents a murine tmst2-receptor polypeptide clone of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 133 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; SEQ ID NO 6; 57pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                  121
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                                                                                      Toyoshima
                                                                                                                                                                                                                                                                                                                                                                                                                                             DRKCQCRTGLYYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MFGFFCSIVSSISRWFIWRRILLLLLLLLLLLNIPLQVKFAMLELHSFKCFAGEYWSKDVCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DRKCOCRTGLYYY
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                                                                                                                                                                                                                                                                                        differentiation; osteoblast cell; bone growth; bone sarcoma.
                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                             97JP-00099653.
                                                                                                                                                  98WO-JP001511.
                                                                                                                                                                                                              /note= "mature protein"
                                                                                                                                                                                                                                  /note= "signal peptide"
                                                                                                                                                                                                                                                        Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                         protein
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Pred. No. 5.9e-54;
                                                                                                                                                                                                                                                                                                                          7F4.
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              in osteoblast cells -
for screening substance
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                substances
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               for
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Claim 1; Page 29-31; Sipp; Japanese

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RESULT 10
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Matches 142;
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         Novel ymkz5-receptor polypeptide useful for treating diseases such as tumor, cancer, AIDS, anemia, autoimmune diseases, cachexia, leprosy, leukemia, hepatitis, multiple sclerosis.
                                                                     WPI; 2004-008943/01.
N-PSDB; ADF57550.
                                                                                                          Zhang K;
                                                                                                                                                                                           11-JUL-2002; 2002US-00193616
                                                                                                                                                                                                                                                                                                              Transmembrane decoy receptor; ymkz5; tumour necrosis factor; cancer; acquired immune deficiency syndrome; AIDS; anaemia; autoimmune disease; cachexia; leprosy; leukaemia; hepatitis;
                                                                                                                                                                                                                                               US2003096355-A1
                                                                                                                                                                                                                                                                       Mus musculus
                                                                                                                                                                                                                                                                                                      multiple sclerosis; myocardial
                                                                                                                                   (/NAHZ)
                                                                                                                                                                                                                                                                                                                                                                           Mouse ymkz5
                                                                                                                                                                                                                                                                                                                                                                                                                             ADF57551;
                                                                                                                                                                                                                                                                                                                                                                                                                                              ADF57551 standard; protein; 176 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence represents a protein designated 7F4. This protein is capable of inducing differentiation in osteoblast cells. The protein may be used to screen compounds for the ability to bind to it, for use as differentiation inducing activity. To be a differentiation inducing activity. Compounds so identified, as well as the protein itself, DNA encoding it, and antibodies to it, may be used in the treatment of diseases of bone growth and osteoblast differentiation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TFVKAPCKI PHTQGQCEKCHPGT FTGKONGLHDCELCSTCDKDQNMVADCSATSDRKCEC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TFVKAPCEIPHTQGQCEKCHPGTFTEKDNYLDACILCSTCDKDQEMVADCSATSDRKCQC 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SLVSSLSRWFLWRRLLLLLLLLLLNLPLQVKFAMLELHSFKCPAGEYWSKDVCCKNCSAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SHVSSLSHWP-----LLLLLLNLFLPVTFAMPESYSFNCPDGEYQSNDVCCKTCPSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   176 AA;
                                                                                                                                                       99US-0143137P.
2000US-00611989.
                                                                                                                                                                                                                                                                                                                                                                       receptor.
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                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           68.7%;
75.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8;
                                                                                                                                                                                                                                                                                                ischaemia; obesity; gene therapy; mouse
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                                                                                                                                                                                                                                                                                                                                    TNF; tumour;
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Claim 13; SEQ ID NO

8

57pp; English

The present invention relates to a transgenic non-human animal having the expression of 7F4 gene artificially modified. The transgenic animals are a disease model for bone and glycolipid metabolism disorders. Substances identified by the screening method are agents for the prevention and

Claim 3; SEQ ID NO 2; 44pp;

Japanese.

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នួន
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches 142;
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Best Local
                                                                                                                                               Kake T,
                                                                                                                                                                                                            10-SEP-2003; 2003WO-JP011545.
                                                                            screening
                                                                                     Transgenic non-human animal
                                                                                                                                                                                       17-SEP-2002; 2002JP-00270321
                                                                                                                                                                                                                                                           WO2004026026-A1
                                                                                                                                                                                                                                                                                 Mus musculus
                                                                                                                                                                                                                                                                                                  7F4 gene; Osteopathic; Anorectic; Antidiabetic; glycolipid metabolism disorder; osteoporosis; o
                                                                                                                                                                 (CHUS ) CHUGAI SEIYAKU
                                                                                                                                                                                                                                     01-APR-2004
                                                                                                                                                                                                                                                                                                                                          Mouse 7F4
                                                                                                                                                                                                                                                                                                                                                               17-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                       ADM46623;
                                                                                                                                                                                                                                                                                                                                                                                                       ADM46623 standard; protein; 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to transmembrane decoy receptor, ymkz5 belonging to tumour necrosis factor (TNP) receptor supergene family and nucleic acid sequences encoding such receptors. The invention is useful for detecting ymkz5-receptor gene such as tumours or cancers. The sequences of mutated invention are used as medication for a number of diseases such as acquired immune deficiency syndrome (AIDS), anaemia, autoimmune diseases, cachexia, leprosy, leukaemia, hepatitis, multiple sclerosis, myocardial ischaemia, obesity etc. The invention is also useful in gene therapy. The present sequence is mouse ymkz5 receptor protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 176 AA;
                                                                                                           2004-340227/31.
DB; ADM46622.
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                                                                                                                                            Saito H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ഗ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7 SLVSSLSRWFLWRRLLLLLLLLLLLLLLDLQVKFAMLELHSFKCPAGEYWSKDVCCKNCSAG
                                                                           remedies for bone
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                                                                                                                                                                                                                                                                                                                                         protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SHVSSLSHWF-----LLLLLLLLFLPVIFAMPESYSPNCPDGEYQSNDVCCKTCPSG
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                                                                                                                                                                                                                                                                                                                                                             (first
                                                                                                                                           Makishima
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                                                                      with modified expression or glycolipid metabolism
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 746.5;
Pred. No. 1e-5
8; Mismatches
                                                                                                                                                                                                                                                                                               osteoporosis; obesity; diabetes.
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                                                                    of 7F4 gene disorders.
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Best Local S
Matches 402
                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence encodes a protein designated 7F4. This protein is capable of inducing differentiation in osteoblast cells. The protein may be used to screen compounds for the ability to bind to it, for use as ligands, agonists or antagonists and inhibiting or otherwise altering its differentiation inducing activity. Compounds so identified, as well as the protein itself, DNA encoding it, and antibodies to it, may be used in the treatment of diseases of bone growth and osteoblast differentiation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Receptor protein inducing differentiation extracellular region only and can be used treatment of bone growth disorders.
                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1509 BP; 387 A; 403
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kimura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-APR-1998;
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              501
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TGTGTGCAGTTCATCTGTTTCAA
                                                ATGTACCAAGTGTCCCCAAGGAATCCCTGTCCTCCAGGAATGCAACTCCACAGCTAACAC
                                                                               ATGCGAGTGCCAAATAGGTCTTTACTACTATGACCCAAAATTTCCGGAATCATGCCGCCC
                                                                                            ATGCCAGTGCCGAACAGGTCTTTACTACTATGACCCAAAATTTTCCAGAATCGTGCCGCCC
                                                                                                                          CTCCACCTGTGATAAAGACCAGAATATGGTGGCTGACTGTTCTGCCACCAGTGACCGGAA
                                                                                                                                          GAAGTGTCACCCAGGAACATTCACAGGGAAAGATAATGGCCTGCATGATTGTGAACTTTG
                                                                                                                                                                                            GAAGTGTCACCCAGGAACATTCACAGAGAAAGATAATTACCTGGATGCTTGTATACTTTG
                                                                                                                                                                                                                     TCCCTCAGGTACATTTGTCAAGGCGCCCTGCAAAATCCCCCATACTCAAGGACAATGTGA
                                                                                                                                                                                                                                             TTCTGCAGGTACATTTGTCAAGGCGCCCTGCGAAATCCCCCCATACTCAAGGACAATGTGA
                                                                                                                                                                                                                                                                    ATACTCCTTCAACTGTCCCGATGGTGAATACCAGTCTAATGATGTCTGTTGCAAGACCTG
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Pred. No. 3.4e-98;
0; Mismatches 41;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
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CTCCACCTGTGATAAAGATCAGGAAAATGGTGGCCGACTGCTCAGCCACCAGTGACCGGAA
                                                                               GAAGTGTCACCCAGGAACATTCACAGAGAAAGATAATTACCTGGATGCTTGTATACTTTG
                                                                                                                                                                                                                                                   ACACTCCTTCAAATGTCCCGCTGGTGAAATACTGGTCTAAAGACGTCTGTTGCAAGAACTG
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                                            GAAGTGTCACCCAGGAACATTCACAGGGAAAAGATAATGGCCTGCATGATTGTGAACTTTG
                                                                                                                                                          TTCTGCAGGTACATTTGTCAAGGCGCCCTGCGAAATCCCCCCATACTCAAGGACAATGTGA
                                                                                                                                                                                                                         ATACTCCTTCAACTGTCCCGATGGTGAATACCAGTCTAATGATGTCTGTTGCAAGACCTG
                                                                                                                                                                                                                                                                                                                GTTCCTCTTGCTGCTGCTGAATCTGTTCTTGCCGGTAATATTTGCTATGCCTGAATC
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                                                                                                                                   TCCCTCAGGTACATTTGTCAAGGCGCCCTGCAAAATCCCCCATACTCAAGGACAATGTGA
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remedies for bone or glycolipid
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d metabolism d
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/product= "7F4"
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Pred. No. 3.4
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disorders.
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RESULT 12
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                                                                                                                                                                                                                                  The present invention relates to a transgenic non-human animal having the expression of 7F4 gene artificially modified. The transgenic animals are a disease model for bone and glycolipid metabolism disorders. Substances identified by the screening method are agents for the prevention and treatment of diseases including osteoporosis, obesity and diabetes. The present sequence represents the modified mouse 7F4 protein.
                                                                                                                                                                                                                                                                                                                                                                       Transgenic non-human animal with modified screening remedies for bone or glycolipid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kake T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-SEP-2002;
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                                                                                                                                                                                                           Sequence 176 AA;
                                                                                                                                                                                                                                                                                                                                              Claim 3; SEQ ID NO 2; 44pp; Japanese.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TFVKAPCKI PHTQGQCEKCHPGTFTGKDNGLHDCELCSTCDKDQNMVADCSATSDRKCEC
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                                            TFVKAPCEIPHTOGOCEKCHPGTFTEKDNYLDACILCSTCDKDQEMVADCSATSDRKCQC
RTGLYYYDPKFPESCRPCTKCPQGIPVLQECNSTANTVCSSSVS 170
                               TFVKAPCKIPHTQGQCEKCHPGTPTGKDNGLHDCELCSTCDKDQNMVADCSATSDRKCEC
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                                                                                                                                                 Score 705; DB 8; ]
Pred. No. 2.1e-50;
5; Mismatches 20;
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                                                                                                                                                                                                                                                                                                                                      Sequence 148 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 3; SEQ ID NO 3; 44pp; Japanese.
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                                                                         ABG09344
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                                                                                                                                                           159
                                                                                                                                                                                                                                                                                      112;
                                                                                                                                                                                                                                                              39
                                                                                                                                                                                   13
                                                                                                                                                                                                          99 ACILCSTCDKDQEMVADCSATSDRKCQCRTGLYYYDPKFPESCRECTKCPQGIPVLQECN
                                                                                                                                                                                                                                     ٠,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CHUGAI SEIYAKU KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Saito H,
                                                                                                                                                                                   DCBLCSTCDKDQNWVADCSATSDRKCECQIGLYYYDPKFPBSCRPCTKCPQGIPVLQBCN
                                                                                                                                                                                                                                    AMPESYSPNCPDGEYQSNDVCCKTCPSGTPVKAPCKIPHTQGQCEKCHPGTFTGKDNGLH
                                                                                                                                                                                                                                                    AMLELHSFKCPAGEYMSKDVCCKNCSAGTFVKAPCEIPHTQGQCEKCHPGTFTEKDNYLD
                                                                          standard;
                                                                                                                                                           STANTYCSSSVS 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein
                                                                                                                                     STANTVCSSSVS 132
                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Makishima
                                                                         protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entry)
                                                                                                                                                                                                                                                                                                  63.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                148
                                                                                                                                                                                                                                                                                       5
                                                                           380
                                                                                                                                                                                                                                                                                      Score 641; DB 8;
Pred. No. 3.3e-45;
5; Mismatches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  osteoporosis; obesity; diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                B
                                                                                                                                                                                                                                                                                                                Length 148;
                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tor
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Novel human diagnostic protein #9335.

13-FEB-2002 ABG09344;

(first entry)

158 60

98 0

120